## **Amendment to the Claims**

Claims 1–4. Canceled

- 5. (Currently amended): The An isolated nucleic acid molecule of claim 1, comprising a nucleotide sequence which encodes a polypeptide having an amino acid sequence of SEQ ID NO: 12 or an amino acid sequence having at least 40% sequence identity thereto, to an amino acid sequence selected from the group consisting of SEQ ID NOS: 2, 4, 6, 8, 10 and 12 wherein said polypeptide is a transmembrane protein which has 2,5-diketo-gluconate (2,5-DKG) permease activity.
- 6. (Currently amended): The isolated nucleic acid molecule of elaim 1 claim 5, comprising a nucleotide sequence which encodes a polypeptide having at least 80% sequence identity to an amino acid sequence selected from the group consisting of SEQ ID NOS: 2, 4, 6, 8, 10 and 12 the amino acid sequence of SEQ ID NO: 12.
- 7. (Currently amended): The isolated nucleic acid molecule of elaim 1 claim 5, which encodes a polypeptide having an amino acid sequence selected from the group consisting of SEQ ID NOS: 2, 4, 6, 8, 10 and 12 the amino acid sequence of SEQ ID NO: 12.

Claims 8 – 10. Canceled

- 11. (Currently amended): The An isolated nucleic acid molecule of claim 1 comprising a polynucleotide which encodes a polypeptide having an amino acid sequence of SEQ ID NO.

  12 or an amino acid sequence having at least 40% sequence identity thereto, wherein said polypeptide has 2,5-diketo-gluconate (2,5-DKG) permease activity, and wherein the polynucleotide is operatively linked to a promoter of gene expression.
- 12. (Original): The isolated nucleic acid molecule of claim 11, wherein said promoter is a *lac* promoter.

13. (Original): A vector comprising the isolated nucleic acid molecule of claim 11.

14. (Original): The vector of claim 13, comprising a spectromycin resistance gene.

15. (Original): A bacterial cell, comprising the vector of claim 13.

16. (Currently amended): The bacterial cell of claim 15, wherein said isolated nucleic acid molecule comprises a nucleotide sequence which encodes a polypeptide having an amino acid sequence at least 80% identical to an the amino acid sequence selected from the group eonsisting of SEQ ID NO: 2, 4, 6, 8, 10 and 12 of SEQ ID NO: 12.

Claims 17 - 19. Canceled.

20. (Original): The bacterial cell of claim 15, which is of the genus *Klebsiella*.

21. (Original): The bacteria cell of claim 15, which is deficient in endogenous 2,5-DKG activity.

22. (Currently amended): The bacterial cell of claim 21, <u>further comprising</u> an isolated nucleic acid molecule encoding a polypeptide having <u>2-keto reductase activity and</u> at least 80% <u>sequence</u> identity to SEQ ID NO: 14 and <u>2-keto reductase activity</u>.

23. (Currently amended): The bacterial cell of claim 21, <u>further comprising an isolated</u> nucleic acid molecule encoding a polypeptide having <u>5-keto reductase activity and</u> at least 80% <u>sequence</u> identity to SEQ ID NO: 16 and <u>5-keto reductase activity</u>.

24. (Original): The bacterial cell of claim 15, which is of the genus *Pantoea*.

25. (Currently amended): The bacterial cell of claim 15, which expresses an enzyme that catalyzes the conversion of 2,5-DKG to 2-KLG 2-keto-L-gulonic acid (2-KLG).

26. (Original): The bacterial cell of claim 25, which expresses enzymes that catalyze the conversion of glucose to 2,5-DKG.

27. (Original): The bacterial cell of claim 26, which is deficient in endogenous 2-keto-reductase activity.

Claims 28 – 35. Canceled

36. (Currently amended): A method of using the isolated nucleic acid molecule of claim 15 to enhance 2-KLG 2-keto-L-gulonic acid (2-KLG) production, comprising expressing the polypeptide encoded by said a) introducing the isolated nucleic acid molecule of claim 5 into in a bacterial cell which expresses an enzyme that catalyzes the conversion of 2,5-DKG to 2-KLG, b) allowing expression of the polypeptide encoded by said nucleic acid molecule and c) culturing the bacterial cell under suitable conditions to produce 2-KLG.

37. (Original): The method of claim 36, wherein said bacterial cell further expresses enzymes that catalyze the conversion of glucose to 2,5-DKG.

38. (Original): The method of claim 37, wherein said bacterial cell is deficient in endogenous 2-keto reductase activity.

39. (Original): The method of claim 36, wherein said bacterial cell is of the genus *Pantoea*.

40. (Original): The method of claim 36, further comprising converting said 2-KLG to ascorbic acid.

(Div of 09/922,501) Page 6

Claims 41 - 48. Canceled

49. (New): The bacterial cell of claim 15, which is an *E. coli* cell.

50. (New): The method of claim 36, wherein the nucleic acid molecule encodes a polypeptide having at least 80% sequence identity to SEQ ID NO: 12.

51. (New): The method of claim 36, wherein the nucleic acid molecule has the sequence of SEQ ID NO: 11 or a sequence having at least 95% sequence identity thereto.

52. (New): A method for increasing the transport of 2, 5-DKG across a cell membrane into a bacterial host cell comprising a) introducing the nucleic acid molecule of claim 5 having 2,5-DKG permease activity into a bacterial host cell, b) allowing expression of the 2,5-DKG permease and c) culturing the bacterial host cell under suitable conditions for the transport of 2,5-DKG into the bacterial host cell.

53. (New): The method according to claim 52, wherein the bacterial host cell is an *E. coli*, *Pantoea* or *Klebsiella* host cell.

54. (New): The method according to claim 52, wherein the nucleic acid molecule encodes a polypeptide having at least 80 % sequence identity to SEQ ID NO: 12.

55. (New): The method according to claim 52, wherein the nucleic acid molecule has the sequence of SEQ ID NO: 11 or a sequence having at least 95% sequence identity thereto.

56. (New): An isolated oligonucleotide comprising at least 20 contiguous nucleotides of the nucleotide sequence of SEQ ID NO: 11, wherein said oligonucleotide is used as a probe and hybridizes under stringent hybridization conditions to a nucleic acid that encodes a polypeptide having 2,5-diketo-D-gluconic acid permease activity.